

10/539093

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/539,093
Source: PCP
Date Processed by STIC: 4/11/06

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PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/539,093

DATE: 04/11/2006

TIME: 12:33:16

Input Set : E:\2394 Pr.ST25.txt
 Output Set: N:\CRF4\04112006\J539093.raw

3 <110> APPLICANT: Alcon, Inc.
 4 Yanni, John M.
 5 Gamache, Daniel A.
 6 Miller, Steven T.
 8 <120> TITLE OF INVENTION: Treatment of Dry Eye in Postmenopausal Women by Restoring
 9 15-lipoxygenase Activity to Ocular Surface Cells
 11 <130> FILE REFERENCE: 2394 US
 C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/539,093
 C--> 14 <141> CURRENT FILING DATE: 2005-06-15
 16 <150> PRIOR APPLICATION NUMBER: US 60/435,988
 17 <151> PRIOR FILING DATE: 2002-12-20
 19 <160> NUMBER OF SEQ ID NOS: 10
 21 <170> SOFTWARE: PatentIn version 3.3
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 24 <211> LENGTH: 2671
 25 <212> TYPE: DNA
 26 <213> ORGANISM: homo sapiens
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93	cggaatgcaa	agctggacat	gcccctacgag	tacctgcggc	ccagcgtgtt	ggaaaacagt	1980
95	gtggccatct	aagcgtcgcc	accctttggt	tatttcagcc	cccatcaccc	aagccacaag	2040
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99	cctccctaga	ggggcacctt	ttcatggct	ctgcacccag	tgaacacatt	ttactctaga	2160
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121 <211> LENGTH: 661

122 <212> TYPE: PRT

123 <213> ORGANISM: homo sapiens

125 <400> SEQUENCE: 2

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132					20				25						30	
135	Ala	Ala	Leu	Gly	Lys	Arg	Leu	Trp	Pro	Ala	Arg	Gly	Glu	Thr	Glu	Leu
136					35				40						45	
139	Lys	Val	Glu	Val	Pro	Glu	Tyr	Leu	Gly	Pro	Leu	Leu	Phe	Val	Lys	Leu
140					50				55						60	
143	Arg	Lys	Arg	His	Leu	Leu	Lys	Asp	Asp	Ala	Trp	Phe	Cys	Asn	Trp	Ile
144					65				70						80	
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148					85				90						95	
151	Arg	Trp	Val	Glu	Gly	Asn	Gly	Val	Leu	Ser	Leu	Pro	Glu	Gly	Thr	Gly
152					100				105						110	
155	Arg	Thr	Val	Gly	Glu	Asp	Pro	Gln	Gly	Leu	Phe	Gln	Lys	His	Arg	Glu
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159	Glu	Glu	Leu	Glu	Glu	Arg	Arg	Lys	Leu	Tyr	Arg	Trp	Gly	Asn	Trp	Lys
160					130				135						140	
163	Asp	Gly	Leu	Ile	Leu	Asn	Met	Ala	Gly	Ala	Lys	Leu	Tyr	Asp	Leu	Pro
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175 Leu Thr Cys Trp Lys Asp Leu Asp Asp Phe Asn Arg Ile Phe Trp Cys
 176 195 200 205
 179 Gly Gln Ser Lys Leu Ala Glu Arg Val Arg Asp Ser Trp Lys Glu Asp
 180 210 215 220
 183 Ala Leu Phe Gly Tyr Gln Phe Leu Asn Gly Ala Asn Pro Val Val Leu
 184 225 230 235 240
 187 Arg Arg Ser Ala His Leu Pro Ala Arg Leu Val Phe Pro Pro Gly Met
 188 245 250 255
 191 Glu Glu Leu Gln Ala Gln Leu Glu Lys Glu Leu Glu Gly Gly Thr Leu
 192 260 265 270
 195 Phe Glu Ala Asp Phe Ser Leu Leu Asp Gly Ile Lys Ala Asn Val Ile
 196 275 280 285
 199 Leu Cys Ser Gln Gln His Leu Ala Ala Pro Leu Val Met Leu Lys Leu
 200 290 295 300
 203 Gln Pro Asp Gly Lys Leu Leu Pro Met Val Ile Gln Leu Gln Leu Pro
 204 305 310 315 320
 207 Arg Thr Gly Ser Pro Pro Pro Pro Leu Phe Leu Pro Thr Asp Pro Pro
 208 325 330 335
 211 Met Ala Trp Leu Leu Ala Lys Cys Trp Val Arg Ser Ser Asp Phe Gln
 212 340 345 350
 215 Leu His Glu Leu Gln Ser His Leu Leu Arg Gly His Leu Met Ala Glu
 216 355 360 365
 219 Val Ile Val Val Ala Thr Met Arg Cys Leu Pro Ser Ile His Pro Ile
 220 370 375 380
 223 Phe Lys Leu Ile Ile Pro His Leu Arg Tyr Thr Leu Glu Ile Asn Val
 224 385 390 395 400
 227 Arg Ala Arg Thr Gly Leu Val Ser Asp Met Gly Ile Phe Asp Gln Ile
 228 405 410 415
 231 Met Ser Thr Gly Gly Gly His Val Gln Leu Leu Lys Gln Ala Gly
 232 420 425 430
 235 Ala Phe Leu Thr Tyr Ser Ser Phe Cys Pro Pro Asp Asp Leu Ala Asp
 236 435 440 445
 239 Arg Gly Leu Leu Gly Val Lys Ser Ser Phe Tyr Ala Gln Asp Ala Leu
 240 450 455 460
 243 Arg Leu Trp Glu Ile Ile Tyr Arg Tyr Val Glu Gly Ile Val Ser Leu
 244 465 470 475 480
 247 His Tyr Lys Thr Asp Val Ala Val Lys Asp Asp Pro Glu Leu Gln Thr
 248 485 490 495
 251 Trp Cys Arg Glu Ile Thr Glu Ile Gly Leu Gln Gly Ala Gln Asp Arg
 252 500 505 510
 255 Gly Phe Pro Val Ser Leu Gln Ala Arg Asp Gln Val Cys His Phe Val
 256 515 520 525
 259 Thr Met Cys Ile Phe Thr Cys Thr Gly Gln His Ala Ser Val His Leu
 260 530 535 540
 263 Gly Gln Leu Asp Trp Tyr Ser Trp Val Pro Asn Ala Pro Cys Thr Met
 264 545 550 555 560
 267 Arg Leu Pro Pro Pro Thr Thr Lys Asp Ala Thr Leu Glu Thr Val Met
 268 565 570 575
 271 Ala Thr Leu Pro Asn Phe His Gln Ala Ser Leu Gln Met Ser Ile Thr

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276 595 600 605
279 Glu Glu Glu Tyr Phe Ser Gly Pro Glu Pro Lys Ala Val Leu Lys Lys
280 610 615 620
283 Phe Arg Glu Glu Leu Ala Ala Leu Asp Lys Glu Ile Glu Ile Arg Asn
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297 <212> TYPE: DNA
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305 ccaacggagat aaccccttag tacatctggc ccatctggc aaggagtca gcgcggjt
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412 <212> TYPE: PRT

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426 35 40 45
429 Ala Glu Glu Asp Phe Glu Val Thr Leu Pro Gln Asp Val Gly Thr Val
430 50 55 60
433 Leu Met Leu Arg Val His Lys Ala Pro Pro Glu Val Ser Leu Pro Leu
434 65 70 75 80
437 Met Ser Phe Arg Ser Asp Ala Trp Phe Cys Arg Trp Phe Glu Leu Glu
438 85 90 95
441 Trp Leu Pro Gly Ala Ala Leu His Phe Pro Cys Tyr Gln Trp Leu Glu
442 100 105 110
445 Gly Ala Gly Glu Leu Val Leu Arg Glu Gly Ala Ala Lys Val Ser Trp
446 115 120 125
449 Gln Asp His His Pro Thr Leu Gln Asp Gln Arg Gln Lys Glu Leu Glu
450 130 135 140
453 Ser Arg Gln Lys Met Tyr Ser Trp Lys Thr Tyr Ile Glu Gly Trp Pro
454 145 150 155 160
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458 165 170 175
461 Tyr Ser Ala Met Lys Asn Ala Lys Leu Phe Phe Lys Ala His Ser Ala

VERIFICATION SUMMARY

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L:13 M:270 C: Current Application Number differs, Replaced Current Application Number

L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date